

AMENDMENTS TO THE SPECIFICATION

Please amend the paragraph spanning pages 54-55 as follows:

DNA and protein sequence analysis and homology search by BLASTN, TBLASTN, and BLASTP were carried out using programs available from the NCBI website (~~http~~ [://www.ncbi.nlm.nih.org/](http://www.ncbi.nlm.nih.org/)). Databases used include those from the NCBI site, the Sanger Genome Centre website (<http://www.sanger.ac.uk/Projects/Microbes>), and the SwissProt website (<http://www.expasy.org/sprot/>). The positions of the LEE PAI as well as the prophages in the EHEC genome were obtained from data generated by the IslandPath program (<http://www.pathogenomics.sfu.ca/islandpath/>) (13).